



3/6

## Replacement Sheet

RECEIVED

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underlined = deleted in targeting construct

[ ] = sequence flanking Neo insert in targeting construct

TECH CENTER 1600/2900

[CAGCGTCAGACGCAGGGCACTGAGAATGTGCGACAGCGCGAACGATGAAGTAGCCAG  
AGGGTCCCTGGAAAATGAGGCCAGGGTCCC] TGCTGCTGCTTGTCTGCTCGCCCT  
GTCCAGGAGCCTGCGGGCAAAGAGTGTGCGTCTCCACCCGTGA [GTGTCACCAGGAGG  
ACGACTTCAGAGTCACCTGCAAGGAGCTCCACCGAACATCCCCAGCCTGCCGCCAGCACCC  
AGACTCT] GAAGCTCATCGAGACTCATCTGAAGACCATAcccAGTCTTGCAATTTCGAGT  
CTGCCAATATTTCCAGGATCTATTATCTATAGATGCAACTCTGCAGCGGCTGGAACCA  
CATTCTTCTACAATTGAGTAAAATGACTCACATAGAAATCCGAACACCAGAAGCTTA  
ACCTATATAGACCCTGATGCCTTGACAGAGCTCCCTGCTCAAGTTCTTGGCATTTC  
AATACTGGACTTAGAATATTCCCTGACTTGACCAAAATTATTCCACGGACATATTCTT  
ATACTTGAAATCACAGACAACCCCTACATGACTTCGGCCCTGAAAACGCATTCCAGGGC  
CTATGCAATGAAACCTGACCCCTGAAACTGTACAACAAATGGATTACTTCAGTCCAAGGA  
CATGCTTCATGGAACAAAGCTGGATGCTGTTACCTAAACAAGAATAAACACTGACA  
GCTATAGACAACGATGCCCTTGGAGGAGTATAACAGTGGACCAACTTGCTAGATGTCT  
TCCACCAGCGTCACTGCCCTTCCAAAGGCCCTGGAGCACCTCAAAGAACTGATCGCA  
AAAGACACCTGGACTCTCAAAAGCTCCGCTGCGTTAACCTCGTCAAAGGAAATCA  
GCTGACCTCTCTTACCCGAGCCACTGCTGCGCTTTAAGAACCAAGAAGAAAATCAGGGGA  
ATCCTGGAGTCCTTGATGTAAATGAGAGCAGTATCCGGAACCTTCGTCAAAGGAAATCA  
GTGAACATCTTGAGGGTCCCCTACAGGAAATATGAAGAACATCCGGTGACAACAGT  
GTTGGGTACAAACAAACTCCAAGTTCCAGGAGAGCCAAAGCAACTCTCACTATTACGTC  
TTCTTGAAGAACAAAGAGGATGAGGTGTTGGCTGGCAAGAGCTCAAAGGAAATC  
GAAGAGACTCTCCAAGCCTCGAGAGCCACTATGACTACACGGTGTGTTGGGACAACGAG  
GACATGGTGTGACCCCCAAGTCGGACGAGTTAACCCCTGTGAAGATATCATGGCTAC  
AGGTTCTGAGAACATGTTGGTGTGGTTGTCAGTCTGCTGGCTCTCCTGGCAATATCTT  
GTCCTGCTCATCTGCTAACCAAGCCACTACAAATTGACCGTGCGCGGTTCTCATGTGC  
AACTTGGCCTTGCAGATTCTGCATGGGGTATACCTGCTTCTCATTGCCTCTGTAGAC  
CTGTACACACACTCTGAGTACTACAACCACGCCATCGACTGGCAGACGGGCCCTGGGTGC  
AACACGGCTGGCTTCACTGTTGCCAGTGAGTTACAGTGTACACACTGACGGTC  
ATCACCCCTGGAGCGATGGTACGCCATCACCTCGCCATGCGCCTGGATAGGAAGATCCGC  
CTCAGGCACCGTACACCACATGGCTGGGGCTGGGTTCTGCTTCTCGCCCTG  
CTCCCGATGGTGGGAATCAGCAGCTATGCCAAGGTAGCAGCATCTGCCCTGCCAATGGACACC  
GACACCCCTCTGCACTCGCATACATTGCTCGTTCTGCTGCTCAATGTTGCTT  
GTTGTCGTCTGTTCTGCTATGTGAAGATCTACATCACGGTCCGAAATCCCCAGTACAAC  
CCTCGAGATAAAGACACCAAGATTGCCAAGAGGGATGGCTGTGATCTCACTGACTTC  
ATGTGCATGGGCCCATCTCCTCTATGCGCTGTCGGCACTTATGAACAAGCCTCTAATC  
ACTGTTACTAACTCCAAAATCTGTTGGTCTCTTCTACCCCTCAACTCCTGTGCCAAT  
CCGTTCTATGCTATTTACCAAGGCCCTCCAGAGGGACGTGTTCATCCTGCTCAGC  
AAGTTGGCATCTGCAAACGCCAGGCCAGGCCTATCAGGGTCAGAGAGTCTGCTT  
AATAGCACTGGTATTCAAGATCCAAAAGATTCCCCAGGACACGAGGAGTCTCCCCAAC  
ATGCAAGATACTTGAACACTGCTGGAAACTCCCAGCTAGCTCCAAAATGCAGGGACAA  
ATCTCAGAACAGAGTATAAGCAAACAGCCTTGTAAAGGAAAGGCTACGCTAGTCACAGTGAG  
ACTTACAAAAGGCTGGTTCTGAACATGCGTCCAGTCCGTGACATGTGAACACATAG  
GTTCATGCAAGGTGATGATTCAAGGGTCAGAGTTCATAGAAAGTATTGCCTC  
(SEQ ID NO:1)

FIGURE 2A